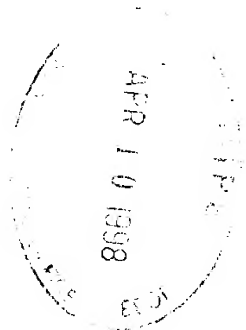


Fig 4

FOR LINE-UP (ref)



PCTED	1	-----	-----	-----	-----	50
BOVGSTA	1	CCCCCCCC	CCCCCCCC	CCCCCCCC	CCCCCCCC	50
MUSCULTNS	1	-----	-----	-----	-----	50
PCTED	51	-----	-----	-----	-----	100
BOVGSTA	51	CCCCCCCC	CCCCCCCC	CCCCCCCC	CCCCCCCC	100
MUSCULTNS	51	-----	-----	-----	-----	100
PCTED	101	-----	-----	-----	-----	150
BOVGSTA	101	CCCCCCCC	CCCCCCCC	CCCCCCCC	CCCCCCCC	150
MUSCULTNS	101	-----	-----	-----	-----	150
PCTED	151	-----	-----	-----	-----	200
BOVGSTA	151	CCCCCCCC	CCCCCCCC	CCCCCCCC	CCCCCCCC	200
MUSCULTNS	151	-----	-----	-----	-----	200
PCTED	201	-----	-----	-----	-----	250
BOVGSTA	201	CCCCCCCC	CCCCCCCC	CCCCCCCC	CCCCCCCC	250
MUSCULTNS	201	-----	-----	-----	-----	250
PCTED	251	-----	-----	-----	-----	300
BOVGSTA	251	CCCCCCCC	CCCCCCCC	CCCCCCCC	CCCCCCCC	300
MUSCULTNS	251	-----	-----	-----	-----	300
PCTED	301	-----	-----	-----	-----	350
BOVGSTA	301	CCCCCCCC	CCCCCCCC	CCCCCCCC	CCCCCCCC	350
MUSCULTNS	301	-----	-----	-----	-----	350
PCTED	351	-----	-----	-----	-----	400
BOVGSTA	351	CCCCCCCC	CCCCCCCC	CCCCCCCC	CCCCCCCC	400
MUSCULTNS	351	-----	-----	-----	-----	400
PCTED	401	-----	-----	-----	-----	450
BOVGSTA	401	CCCCCCCC	CCCCCCCC	CCCCCCCC	CCCCCCCC	450
MUSCULTNS	401	-----	-----	-----	-----	450
PCTED	451	-----	-----	-----	-----	500
BOVGSTA	451	CCCCCCCC	CCCCCCCC	CCCCCCCC	CCCCCCCC	500
MUSCULTNS	451	-----	-----	-----	-----	500
PCTED	501	-----	-----	-----	-----	550
BOVGSTA	501	CCCCCCCC	CCCCCCCC	CCCCCCCC	CCCCCCCC	550
MUSCULTNS	501	-----	-----	-----	-----	550
PCTED	551	-----	-----	-----	-----	600
BOVGSTA	551	CCCCCCCC	CCCCCCCC	CCCCCCCC	CCCCCCCC	600
MUSCULTNS	551	-----	-----	-----	-----	600

Exam 1 Exam 2

Exam 2 Exam 3

Fig 4 (cont) 4-2

			Exon 3   Exon 4	
PCTCD	301	-----	-----	650
BOVGSTA	301	-----	-----	650
MUSCULYTHS	301	CAAGTCAGAA	ACAGCTCTTC	650
		CAAG		
PCTCD	651	TAATCATCT	CAAGCAGAA	700
BOVGSTA	651	TAATCATCT	CAAGCAGAA	700
MUSCULYTHS	651	TAATCATCT	CAAGCAGAA	700
			Exon 4   Exon 5	
PCTCD	701	GTAAATGCTT	TCTTTTGGCA	750
BOVGSTA	701	GTAAATGCTT	TCTTTTGGCA	750
MUSCULYTHS	701	GTAAATGCTT	TCTTTTGGCA	750
			Exon 5   Exon 6	
PCTCD	751	CTGCATATAC	CACTCAAAA	800
BOVGSTA	751	CTGCATATAC	CACTCAAAA	800
MUSCULYTHS	751	CTGCATATAT	CACTCAAAA	800
			Exon 6   Exon 7	
PCTCD	801	CTCTCTCTG	TTTCTGGAG	850
BOVGSTA	801	CTCTCTCTG	TTTCTGGAG	850
MUSCULYTHS	801	CTCTCTCTG	TTTCTGGAG	850
PCTCD	851	CAACAGAGAC	ACCTATAGC	900
BOVGSTA	851	CAACAGAGAC	ACCTATAGC	900
MUSCULYTHS	851	CAACAGAGAC	ACCTATAGC	900
			Exon 7   Exon 8	
PCTCD	901	CAAGAGAGAC	CTTCTCTAG	950
BOVGSTA	901	CAAGAGAGAC	CTTCTCTAG	950
MUSCULYTHS	901	CAAGAGAGAC	CTTCTCTAG	950
PCTCD	951	ACCTCTGAC	CAATACCA	1000
BOVGSTA	951	ACCTCTGAC	CAATACCA	1000
MUSCULYTHS	951	ACCTCTGAC	CAATACCA	1000
PCTCD	1001	TACACACAG	CTCTCTTACA	1050
BOVGSTA	1001	TACACACAG	CTCTCTTACA	1050
MUSCULYTHS	1001	TACACACAG	CTCTCTTACA	1050
			Exon 8   Exon 9	
PCTCD	1051	CTCTCTGAC	CTTTTCTG	1100
BOVGSTA	1051	CTCTCTGAC	CTTTTCTG	1100
MUSCULYTHS	1051	CTCTCTGAC	CTTTTCTG	1100
PCTCD	1101	AGTTCTTAAT	ATCTCTTAA	1150
BOVGSTA	1101	AGTTCTTAAT	ATCTCTTAA	1150
MUSCULYTHS	1101	AGTTCTTAAT	ATCTCTTAA	1150
PCTCD	1151	TTTACATCA	TCTCTGATCA	1200
BOVGSTA	1151	TTTACATCA	TCTCTGATCA	1200
MUSCULYTHS	1151	TTTACATCA	TCTCTGATCA	1200
PCTCD	1201	TCTCTCTG	TCTTTTAA	1250
BOVGSTA	1201	TCTCTCTG	TCTTTTAA	1250
MUSCULYTHS	1201	TCTCTCTG	TCTTTTAA	1250

Fig 4(cont) 4-3

POTCD	1151	AACACATCAG	CATCATCCGC	ATCAACACCA	TCCGGGACCA	CATCTCGCC	1200
BOVGSTA	1151	AGCACATCAG	CATCATCCGC	ATCAACACCA	TCCGGGACCA	CATCTCGCC	1200
MUSGLYTHS	1151	AGCATTATCAG	CATCATCCGC	ATCAACACCA	TCCGGGACCA	CATCTCGCC	1200
POTCD	1201	CACATCCAGC	AGCAGCTGCA	CTTCTCTCTC	TCCATCCAGC	TCCATCAAGT	1250
BOVGSTA	1201	CACATCCAGC	ATCAGCTTCA	CTTCTCTCTC	TCCATCCAGC	TCCATCAAGT	1250
MUSGLYTHS	1201	CACATCCAGC	AGCAGCTGCA	CTTCTCTCTC	TCCATCCAGC	TCCATCAAGT	1250
POTCD	1351	CTTCCAAAC	AAGTTTGGCC	TCCAGACTCT	GGCCCACTCC	GTCCCTCACC	1400
BOVGSTA	1351	CTTCCAAAC	AAGTTTGGCC	TCCAGACTCT	GGCCCACTCC	GTCCCTCACC	1400
MUSGLYTHS	1351	CTTCCAAAC	AAGTTTGGCC	TCCAAACTCT	GGCCCACTCC	GTACCAACCC	1400
POTCD	1401	TACAGGCGTC	GTGCTACAC	GCAGATCCCA	ATGACTTCAC	GTACCAACCC	1450
BOVGSTA	1401	TACAGGCGTC	GTGCTACAC	GCAGATCCCA	ATGACTTCAC	GTACCAACCC	1450
MUSGLYTHS	1401	TCCAGGCGTC	GTGCTACAC	GCAGCTCCCA	AGACTTCAC	GTATCAACCC	1450
POTCD	1451	GGCAAGCAGT	GGCAAGCTTA	CATTGGCTTT	GGCAAGCGCC	ATTCTTATTA	1500
BOVGSTA	1451	GGCAAGCAGT	GTGCAAGCTA	CATTGGCTTT	GGCAAGCGCC	ATTCTTATTA	1500
MUSGLYTHS	1451	GGCAAGCTCT	GGCCCGCTTA	CATTGGCTTT	GGCAAGCGCC	ATTCTTATTA	1500
POTCD	1501	GCAGCGCGCC	ATTCTTGGCC	GAACAGCTAC	TCAGCTTCTA	AACATCACTC	1550
BOVGSTA	1501	GCAGCGCGCC	ATTCTTGGCC	GAACAGCTAC	TCAGCTTCTT	AACATCACTC	1550
MUSGLYTHS	1501	GCAGCGCGCC	ATTCTTGGAC	GAACCGCTAC	TCAGCTTCTC	AACATCACTA	1550
POTCD	1551	AGGAGTCTTT	CAAGCGAATC	CTCCAGGACA	AGCAAAATCA	CATAGAACCC	1600
BOVGSTA	1551	AGCAATCTTT	CAAGCGAATC	CTCCAGGACA	AGCAAAATCA	CATAGAACCC	1600
MUSGLYTHS	1551	GGGAGTCTTT	TAAGCGGATC	CTCCAGGACA	AGCAAAATCA	CATAGAACCC	1600
POTCD	1601	CAGTGGCATC	ATCAAGGCA	TCTAAACAC	TATTCTCTTC	TCAGCAACCC	1650
BOVGSTA	1601	CAATGGCATC	ATCAAGGCA	TCTAAACAC	TATTCTCTTC	TCAGCAACCC	1650
MUSGLYTHS	1601	CAGTGGCATC	ATCAAGGCA	CTTCAACAA	TACTCTCTTT	TCAGCAACCC	1650
POTCD	1651	CACAAATATC	TTATCCCGAC	AATATCTCTC	GCATTATCAT	ATAGCGCTAC	1700
BOVGSTA	1651	TACTAAATATC	TTATCCCGCG	AATATCTCTC	GCATTATCAC	ATAGCGCTAC	1700
MUSGLYTHS	1651	CACAAATATC	CTATCTCCAC	AGTATCTCTC	GCATTATCAC	ATAGCGCTCC	1700
POTCD	1701	CTCTCGATAT	TAGCATCTTC	AACATAGCTT	GGCAGAAAA	AGACTATTAAT	1750
BOVGSTA	1701	CTCCCGATAT	TAAGCTCTTC	AACATAGCTT	GGCAGAAAA	AGACTATTAAT	1750
MUSGLYTHS	1701	CTTCAATAT	TAAAGCTCTC	AACATAGCTT	GGCAGAAAA	AGACTATTAAT	1750
POTCD	1751	TTGGTTAGAA	ATAATCTCTC	ACTTTAAATT	GTCCCAACCA	TTTCTCAAT	1800
BOVGSTA	1751	TTGGTTAGAA	ATAATCTCTC	ACTTTAAATT	GTCCCACTAC	ATTCTCAAT	1800
MUSGLYTHS	1751	TTGGTTAGAA	ATAATCTCTC	ACTTTAAATT	GTG-----	---ATCGAAC	1800
POTCD	1801	TTCAACAGT	ATTACTCTCC	CTACTCTCTC	ACAGAACTAC	---CACTTAA	1850
BOVGSTA	1801	TTCAACAGT	ATTACTCTCC	CTACTCTCTC	ACAGAACTAA	---CACTTAA	1850
MUSGLYTHS	1801	TTCAAC--T	ATTACTCTCC	CTAACTCTCTC	AAAGAACTAC	CAAGACTTCA	1850

Stop

Fig. 4(cont) 4 - 4

PGTCD	1051	TTTAACTTT	TAUAAUATA	CTAACAAA-	----TACCAA	CACAGTAA-G	1200
BOVGSTA	1051	TTTAACTTT	AAAAAAATA	CTAACAAA-	----GACCAA	CACAGCAA-A	1200
MUSGLYTHS	1051	TTTCAACTTT	TAUAAUAA-A	CAATCAUAA	CAUAACTTAC	TACCATGCCA	1200
PGTCD	1201	TACATATTAT	TTTCTCTTGC	AACTTTGAGC	CTTCTCAUAT	GCCAGUATCA	1250
BOVGSTA	1201	TACATATTAT	TTTCTCTTGT	AACTTTGAGC	CTTCTCAUAT	GCCAGUATCA	1250
MUSGLYTHS	1201	AACAGATCAT	TTTCTCT-CA	CACCTTTGAGC	CT-CTAATAT	CTCAGUAAU	1250
PGTCD	1251	CTCTCTGC--	--TAATCAGA	TCTAAATTCG	CAGTCAATTC	.....	2000
BOVGSTA	1251	ACCTCTGC--	--TAATCAGA	TCTAAATTCG	CAGTCAATTC	TTACCTATTT	2000
MUSGLYTHS	1251	CTCTATGCCA	AGTAATCAGG	TATAAATTCG	CAATCAATTC	TTATATATTC	2000
PGTCD	2001	.....	.....	.....	.....	.....	2050
BOVGSTA	2001	TTCTCTCTGC	GCCCTCTCTA	TCCATACAGC	ATCATTTCAA	CC.....	2050
MUSGLYTHS	2001	TTCTCTCTGC	CAUAACTTCA	TTCTCAUAT	CAUAAATTAU	TTCTCAUAGC	2050
PGTCD	2051	.....	.....	.....	.....	.....	2100
BOVGSTA	2051	.....	.....	.....	.....	.....	2100
MUSGLYTHS	2051	AAAGCCAT	GCCGAACT	TTCTCTCAGT	CTCTCATACA	ATTACCCACT	2100
PGTCD	2101	.....	.....	.....	.....	.....	2150
BOVGSTA	2101	.....	.....	.....	.....	.....	2150
MUSGLYTHS	2101	GCCAGCTGC	TCAGACAGC	ATTAGGCAAC	ACTCTCTCTT	CTCTCAGACT	2150
PGTCD	2151	.....	.....	.....	.....	.....	2200
BOVGSTA	2151	.....	.....	.....	.....	.....	2200
MUSGLYTHS	2151	TCAGCTCTC	CTCTCTCTT	GCTCAATTAU	CTCTCTCTC	ATCCAGATTC	2200
PGTCD	2201	.....	.....	.....	.....	.....	2250
BOVGSTA	2201	.....	.....	.....	.....	.....	2250
MUSGLYTHS	2201	TAUAGCAGC	CAGAGCTT	TTCCAGCTAC	AGACTTTAA	TAGCATGCC	2250
PGTCD	2251	.....	.....	.....	.....	.....	2300
BOVGSTA	2251	.....	.....	.....	.....	.....	2300
MUSGLYTHS	2251	AACTCTACA	TCATCTCTA	AACTCTCAT	GCTCTAGCA	CCCTCTCAGT	2300
PGTCD	2301	.....	.....	.....	.....	.....	2350
BOVGSTA	2301	.....	.....	.....	.....	.....	2350
MUSGLYTHS	2301	CCAGGCTCTA	CTAGCTATT	CTTCTCTCTC	TCCTCATAAA	CCAGCTCAGC	2350
PGTCD	2351	.....	.....	.....	.....	.....	2400
BOVGSTA	2351	.....	.....	.....	.....	.....	2400
MUSGLYTHS	2351	ACTCTCAATA	CTTACTTTC	TCCTCATCT	TCAGCTCTA	CCAAUATCA	2400
PGTCD	2401	.....	.....	.....	.....	.....	2450
BOVGSTA	2401	.....	.....	.....	.....	.....	2450
MUSGLYTHS	2401	AGCTCTACA	CTTACAGCA	CATAGCACTT	GCTCTCTCTA	ATTCAATCA	2450
PGTCD	2451	.....	.....	.....	.....	.....	2500
BOVGSTA	2451	.....	.....	.....	.....	.....	2500
MUSGLYTHS	2451	CTTACAAACA	CACAGCTT	TTCTCAGCAT	CATCAACAGC	AGCAATTCU	2500

Fig. 4(cont.) 4-5

PCTCD	2501	.....	.....	.....	.....	2550	
BOVGSTA	2501	.....	.....	.....	.....	2550	
MUSGLYTHS	2501	TCGAAAGTGT	GTTCAATTTC	TTTTCCGCAA	ATTGTATCTA	TCGTGTACG	2550
PCTCD	2551	.....	.....	.....	.....	2600	
BOVGSTA	2551	.....	.....	.....	.....	2600	
MUSGLYTHS	2551	TTTGTGTGTT	CAGCGCTGTC	GAGAGCGTCT	CAGTGTATCA	GCGAACATCA	2600
PCTCD	2601	.....	.....	.....	.....	2650	
BOVGSTA	2601	.....	.....	.....	.....	2650	
MUSGLYTHS	2601	GTACCTCAGC	CGACTCAGCA	CGAGCAGCGT	ATTATATCAG	AACACAACTT	2650
PCTCD	2651	.....	.....	.....	.....	2700	
BOVGSTA	2651	.....	.....	.....	.....	2700	
MUSGLYTHS	2651	CTCATCATCA	CGTCTTACCT	ACAACTCTCT	CTGACCTTCC	CAGTCTCTCA	2700
PCTCD	2701	.....	.....	.....	.....	2750	
BOVGSTA	2701	.....	.....	.....	.....	2750	
MUSGLYTHS	2701	CGCCATCTCT	TTCCATCTCT	CGCCGCTCTA	TCGACCATCT	CAGTCTACAC	2750
PCTCD	2751	.....	.....	.....	.....	2800	
BOVGSTA	2751	.....	.....	.....	.....	2800	
MUSGLYTHS	2751	AAAGCCCTCT	GTACATCTAC	CTCATTTCTC	CTGCTCTACT	ACTATCCAGC	2800
PCTCD	2801	.....	.....	.....	.....	2850	
BOVGSTA	2801	.....	.....	.....	.....	2850	
MUSGLYTHS	2801	TGTCACAGCC	AGCCAGCCAG	ATGTACTGCA	CTACATAGCA	ACCCACTTCA	2850
PCTCD	2851	.....	.....	.....	.....	2900	
BOVGSTA	2851	.....	.....	.....	.....	2900	
MUSGLYTHS	2851	TGCGATCTGG	AGCCGCACTC	ACTACACGCC	AGCTGCTCAA	CGTTCTCTTC	2900
PCTCD	2901	.....	.....	.....	.....	2950	
BOVGSTA	2901	.....	.....	.....	.....	2950	
MUSGLYTHS	2901	CGCGCTCTCA	CACTCTGCAG	CAGCCGCTCT	ATTAGTCTTT	CTCACCTTAT	2950
PCTCD	2951	.....	.....	.....	.....	3000	
BOVGSTA	2951	.....	.....	.....	.....	3000	
MUSGLYTHS	2951	CGCTCCGCA	CGCTTTGCA	AGTGTAAAT	GACCTTTCA	CAGCTCTCC	3000
PCTCD	3001	.....	.....	.....	.....	3050	
BOVGSTA	3001	.....	.....	.....	.....	3050	
MUSGLYTHS	3001	CTACACCGCT	TAAAAACAT	AGATATTTC	ACTCTACTCT	CTAACATAG	3050
PCTCD	3051	.....	.....	.....	.....	3100	
BOVGSTA	3051	.....	.....	.....	.....	3100	
MUSGLYTHS	3051	CACAAATCA	GTATCAAAAT	AGCAAGGCA	ATAAATCTTC	CGTTCTCTTC	3100

Fig 5

PDH Line-up (aa):

		Ex4 ↓ Ex5	Ex5 ↓ Ex6				
PQT(Frame 1)	1	HNKCKVLLS	MLVSTAVV	PHYINSPD	SLPHIQSN	PEVG-SSAQR	50
BGT(Frame 1)	1	HNKCKVLLS	MLVSTAVV	PHYINSPD	SLPHIQSN	PEVGSSIQK	50
HGT(Frame 1)	1	HNKCKVLLS	MLVSTAVV	PHYINSPD	SLPHIQSN	PEVGSHRQK	50
		Ex6 ↓ Ex7	Ex7 ↓ Ex8				
PQT(Frame 1)	51	GNWFSWENN	CTHSTHEDD	ATGNKZQRK	EDNAGOLPV	DWNPSEKPS	100
BGT(Frame 1)	51	GNWFSWENN	G---YHEEDG	DTNEKQGRN	ED-SSKLLS	DWNPSEKPS	100
HGT(Frame 1)	51	GNWFSWENN	CTHSTHEDD	EDNAGOLPV	ED-SSKLLS	DWNPSEKPS	100
		Ex8 ↓ Ex9					
PQT(Frame 1)	101	AVTETRWKAP	VWEGTQIRA	VLDNYAKCK	ITVGLTVFV	GRYDHYLED	150
BGT(Frame 1)	101	AVTETRWKAP	VWEGTQIRA	VLDNYAKCK	ITVGLTVFV	GRYDHYLED	150
HGT(Frame 1)	101	AVTETRWKAP	VWEGTQIRA	VLDNYAKCK	ITVGLTVFV	GRYDHYLED	150
PQT(Frame 1)	151	FUSANTYFM	VGRVIFTYM	VDDISAMPLE	ELGPASPKV	FEIKSEKRWQ	200
BGT(Frame 1)	151	FUSANTYFM	VGRVIFTYM	VDDISAMPLE	ELGPASPKV	FEIKSEKRWQ	200
HGT(Frame 1)	151	FUSANTYFM	VGRVIFTYM	VDDISAMPLE	ELGPASPKV	FEIKSEKRWQ	200
PQT(Frame 1)	201	DISMHRMTI	GEHILAHQH	EVDFLFCHDV	DQVFQDNFGV	ETLQGSVAQL	250
BGT(Frame 1)	201	DISMHRMTI	GEHILAHQH	EVDFLFCHDV	DQVFQDNFGV	ETLQGSVAQL	250
HGT(Frame 1)	201	DISMHRMTI	GEHILAHQH	EVDFLFCHDV	DQVFQDNFGV	ETLQGSVAQL	250
PQT(Frame 1)	251	QAHWYKAPD	EFTYERKES	AAIYFFGCGD	FYTHAAIFCG	TPTQVLNITQ	300
BGT(Frame 1)	251	QAHWYKAPD	EFTYERKES	AAIYFFGCGD	FYTHAAIFCG	TPTQVLNITQ	300
HGT(Frame 1)	251	QAHWYKAPD	EFTYERKES	AAIYFFGCGD	FYTHAAIFCG	TPTQVLNITQ	300
PQT(Frame 1)	301	ECFKGILQOX	ENDIEAGHD	ESHNLKYFL	NKPTKILSP	YCADYHIGMS	350
BGT(Frame 1)	301	ECFKGILQOX	ENDIEAGHD	ESHNLKYFL	NKPTKILSP	YCADYHIGMS	350
HGT(Frame 1)	301	ECFKGILQOX	ENDIEAGHD	ESHNLKYFL	NKPTKILSP	YCADYHIGMS	350
PQT(Frame 1)	351	VDIREVKIAN	QKKEYNLVRN	NI*			400
BGT(Frame 1)	351	VDIREVKIAN	QKKEYNLVRN	NV*			400
HGT(Frame 1)	351	VDIREVKIAN	QKKEYNLVRN	NV*			400

